

# **PCT 371 Routing Sheet**

## **APPLICATION**

**IFW DocCode - SEQREQ**

**Index using Current Date**

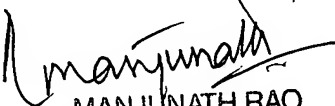
**10/553710**

**TO BE DELIVERED TO:**  
**Tech Center Scanning**

### **Sequence Rule Compliance Review Item**

	CRF, paper copy of sequence listing, and statement that both are same missing
X	CRF contains error(s) according to STIC Report
	CRF damaged or unreadable according to STIC Report
	CRF transferred from prior application is not compliant

**Place an "X" in the appropriate box**

  
MANJUNATH RAO  
SUPERVISORY PATENT EXAMINER  
AV. 1647.

**Comment Sheet**

**APPLICATION SERIAL NUMBER**

**10/553710**

**DOES NOT COMPLY WITH THE  
SEQUENCE RULES. See reasons below.**

**CRF is defective. (see enclosed error  
report)**

**This comment sheet can be used with any of the three  
routing sheets - (SAMPLE) brief reason (e.g., see  
page(s), figure(s), etc. as to why the application is being  
returned. This needs to be type written because it will  
be sent to applicant by PCT, OIPE and PCT/DO/EO.**

**Sample comments should be specific:**

**Page(s) 23, 69 (lines 2 and 23 respectively) contain sequences not found in  
the CRF. See also figures 1A and 4C.**

=====

Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866)  
217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2008; month=6; day=5; hr=15; min=14; sec=13; ms=883; ]

=====

\*\*\*\*\*

Reviewer Comments:

<210> 33

<211> 4

<212> PRT

<213> Artificial Sequence

<220>

<223> exemplary motif

<400> 33

Leu Gly Leu Gly

1

The above <223> response for sequence id# 33 is invalid, please explain  
Artificial. Please correct any other sequences with similar errors.

\*\*\*\*\*

Validated By CRFValidator v 1.0.3

Application No: 10553710

Version No: 1.0

Input Set:

Output Set:

Started: 2008-05-14 15:02:20.012

Finished: 2008-05-14 15:02:21.466

Elapsed: 0 hr(s) 0 min(s) 1 sec(s) 454 ms

Total Warnings: 13

Total Errors: 0

No. of SeqIDs Defined: 37

Actual SeqID Count: 37

Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (1)
W 213	Artificial or Unknown found in <213> in SEQ ID (22)
W 213	Artificial or Unknown found in <213> in SEQ ID (23)
W 213	Artificial or Unknown found in <213> in SEQ ID (26)
W 213	Artificial or Unknown found in <213> in SEQ ID (27)
W 213	Artificial or Unknown found in <213> in SEQ ID (28)
W 213	Artificial or Unknown found in <213> in SEQ ID (29)
W 213	Artificial or Unknown found in <213> in SEQ ID (30)
W 213	Artificial or Unknown found in <213> in SEQ ID (31)
W 213	Artificial or Unknown found in <213> in SEQ ID (33)
W 213	Artificial or Unknown found in <213> in SEQ ID (34)
W 213	Artificial or Unknown found in <213> in SEQ ID (35)
W 213	Artificial or Unknown found in <213> in SEQ ID (36)

# SEQUENCE LISTING

<110> Sah, Dinah Wen-Yee  
Pepinsky, R. Blake  
Rossomando, Anthony

<120> POLYMER-CONJUGATED, GLYCOSYLATED  
NEUBLASTIN

<130> 13751-035W01

<140> 10553710

<141> 2008-05-14

<150> PCT/US04/011745

<151> 2004-04-16

<150> US 60/463,899

<151> 2003-04-18

<160> 37

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 113

<212> PRT

<213> Artificial Sequence

<220>

<223> consensus sequence

<220>

<221> VARIANT

<222> 3

<223> Xaa = Gly or Thr

<220>

<221> VARIANT

<222> 4

<223> Xaa = Pro or Arg

<220>

<221> VARIANT

<222> 5

<223> Xaa = Gly or Ser

<220>

<221> VARIANT

<222> 10, 11

<223> Xaa = Ala or Thr

<220>

<221> VARIANT

<222> 12  
<223> Xaa = Gly or Asp

<220>  
<221> VARIANT  
<222> 26, 33  
<223> Xaa = Arg or Ser

<220>  
<221> VARIANT  
<222> 38, 76  
<223> Xaa = Val or Ile

<220>  
<221> VARIANT  
<222> 53  
<223> Xaa = Pro or Gln

<220>  
<221> VARIANT  
<222> 69  
<223> Xaa = Pro or Ser

<220>  
<221> VARIANT  
<222> 103  
<223> Xaa = Arg or His

<400> 1  
Ala Gly Xaa Xaa Xaa Ser Arg Ala Arg Xaa Xaa Xaa Ala Arg Gly Cys  
1 5 10 15  
Arg Leu Arg Ser Gln Leu Val Pro Val Xaa Ala Leu Gly Leu Gly His  
20 25 30  
Xaa Ser Asp Glu Leu Xaa Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg  
35 40 45  
Arg Ala Arg Ser Xaa His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala  
50 55 60  
Gly Ala Leu Arg Xaa Pro Pro Gly Ser Arg Pro Xaa Ser Gln Pro Cys  
65 70 75 80  
Cys Arg Pro Thr Arg Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser  
85 90 95  
Thr Trp Arg Thr Val Asp Xaa Leu Ser Ala Thr Ala Cys Gly Cys Leu  
100 105 110  
Gly

<210> 2  
<211> 113  
<212> PRT  
<213> Homo sapiens

<400> 2  
Ala Gly Gly Pro Gly Ser Arg Ala Arg Ala Ala Gly Ala Arg Gly Cys  
1 5 10 15  
Arg Leu Arg Ser Gln Leu Val Pro Val Arg Ala Leu Gly Leu Gly His  
20 25 30  
Arg Ser Asp Glu Leu Val Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg

35	40	45
Arg Ala Arg Ser Pro His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala		
50	55	60
Gly Ala Leu Arg Pro Pro Pro Gly Ser Arg Pro Val Ser Gln Pro Cys		
65	70	75
Cys Arg Pro Thr Arg Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser		
85	90	95
Thr Trp Arg Thr Val Asp Arg Leu Ser Ala Thr Ala Cys Gly Cys Leu		
100	105	110
Gly		

<210> 3  
 <211> 113  
 <212> PRT  
 <213> Mus musculus

<400> 3
Ala Gly Thr Arg Ser Ser Arg Ala Arg Thr Thr Asp Ala Arg Gly Cys
1 5 10 15
Arg Leu Arg Ser Gln Leu Val Pro Val Ser Ala Leu Gly Leu Gly His
20 25 30
Ser Ser Asp Glu Leu Ile Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg
35 40 45
Arg Ala Arg Ser Gln His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala
50 55 60
Gly Ala Leu Arg Ser Pro Pro Gly Ser Arg Pro Ile Ser Gln Pro Cys
65 70 75 80
Cys Arg Pro Thr Arg Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser
85 90 95
Thr Trp Arg Thr Val Asp His Leu Ser Ala Thr Ala Cys Gly Cys Leu
100 105 110
Gly

<210> 4  
 <211> 113  
 <212> PRT  
 <213> Rattus norvegicus

<400> 4
Ala Gly Thr Arg Ser Ser Arg Ala Arg Ala Thr Asp Ala Arg Gly Cys
1 5 10 15
Arg Leu Arg Ser Gln Leu Val Pro Val Ser Ala Leu Gly Leu Gly His
20 25 30
Ser Ser Asp Glu Leu Ile Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg
35 40 45
Arg Ala Arg Ser Pro His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala
50 55 60
Gly Ala Leu Arg Ser Pro Pro Gly Ser Arg Pro Ile Ser Gln Pro Cys
65 70 75 80
Cys Arg Pro Thr Arg Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser
85 90 95
Thr Trp Arg Thr Val Asp His Leu Ser Ala Thr Ala Cys Gly Cys Leu
100 105 110
Gly

<210> 5  
 <211> 220  
 <212> PRT  
 <213> Homo sapiens

<400> 5  
 Met Glu Leu Gly Leu Gly Gly Leu Ser Thr Leu Ser His Cys Pro Trp  
 1 5 10 15  
 Pro Arg Arg Gln Pro Ala Leu Trp Pro Thr Leu Ala Ala Leu Ala Leu  
 20 25 30  
 Leu Ser Ser Val Ala Glu Ala Ser Leu Gly Ser Ala Pro Arg Ser Pro  
 35 40 45  
 Ala Pro Arg Glu Gly Pro Pro Val Leu Ala Ser Pro Ala Gly His  
 50 55 60  
 Leu Pro Gly Gly Arg Thr Ala Arg Trp Cys Ser Gly Arg Ala Arg Arg  
 65 70 75 80  
 Pro Pro Pro Gln Pro Ser Arg Pro Ala Pro Pro Pro Pro Ala Pro Pro  
 85 90 95  
 Ser Ala Leu Pro Arg Gly Gly Arg Ala Ala Arg Ala Gly Gly Pro Gly  
 100 105 110  
 Ser Arg Ala Arg Ala Ala Gly Ala Arg Gly Cys Arg Leu Arg Ser Gln  
 115 120 125  
 Leu Val Pro Val Arg Ala Leu Gly Leu Gly His Arg Ser Asp Glu Leu  
 130 135 140  
 Val Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg Arg Ala Arg Ser Pro  
 145 150 155 160  
 His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala Gly Ala Leu Arg Pro  
 165 170 175  
 Pro Pro Gly Ser Arg Pro Val Ser Gln Pro Cys Cys Arg Pro Thr Arg  
 180 185 190  
 Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser Thr Trp Arg Thr Val  
 195 200 205  
 Asp Arg Leu Ser Ala Thr Ala Cys Gly Cys Leu Gly  
 210 215 220

<210> 6  
 <211> 140  
 <212> PRT  
 <213> Homo sapiens

<400> 6  
 Pro Pro Pro Gln Pro Ser Arg Pro Ala Pro Pro Pro Pro Ala Pro Pro  
 1 5 10 15  
 Ser Ala Leu Pro Arg Gly Gly Arg Ala Ala Arg Ala Gly Gly Pro Gly  
 20 25 30  
 Ser Arg Ala Arg Ala Ala Gly Ala Arg Gly Cys Arg Leu Arg Ser Gln  
 35 40 45  
 Leu Val Pro Val Arg Ala Leu Gly Leu Gly His Arg Ser Asp Glu Leu  
 50 55 60  
 Val Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg Arg Ala Arg Ser Pro  
 65 70 75 80  
 His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala Gly Ala Leu Arg Pro  
 85 90 95  
 Pro Pro Gly Ser Arg Pro Val Ser Gln Pro Cys Cys Arg Pro Thr Arg  
 100 105 110  
 Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser Thr Trp Arg Thr Val



115 120 125  
 Asp Arg Leu Ser Ala Thr Ala Cys Gly Cys Leu Gly  
 130 135 140

<210> 7  
 <211> 116  
 <212> PRT  
 <213> Homo sapiens

<400> 7  
 Ala Ala Arg Ala Gly Gly Pro Gly Ser Arg Ala Arg Ala Ala Gly Ala  
 1 5 10 15  
 Arg Gly Cys Arg Leu Arg Ser Gln Leu Val Pro Val Arg Ala Leu Gly  
 20 25 30  
 Leu Gly His Arg Ser Asp Glu Leu Val Arg Phe Arg Phe Cys Ser Gly  
 35 40 45  
 Ser Cys Arg Arg Ala Arg Ser Pro His Asp Leu Ser Leu Ala Ser Leu  
 50 55 60  
 Leu Gly Ala Gly Ala Leu Arg Pro Pro Pro Gly Ser Arg Pro Val Ser  
 65 70 75 80  
 Gln Pro Cys Cys Arg Pro Thr Arg Tyr Glu Ala Val Ser Phe Met Asp  
 85 90 95  
 Val Asn Ser Thr Trp Arg Thr Val Asp Arg Leu Ser Ala Thr Ala Cys  
 100 105 110  
 Gly Cys Leu Gly  
 115

<210> 8  
 <211> 112  
 <212> PRT  
 <213> Homo sapiens

<400> 8  
 Gly Gly Pro Gly Ser Arg Ala Arg Ala Ala Gly Ala Arg Gly Cys Arg  
 1 5 10 15  
 Leu Arg Ser Gln Leu Val Pro Val Arg Ala Leu Gly Leu Gly His Arg  
 20 25 30  
 Ser Asp Glu Leu Val Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg Arg  
 35 40 45  
 Ala Arg Ser Pro His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala Gly  
 50 55 60  
 Ala Leu Arg Pro Pro Pro Gly Ser Arg Pro Val Ser Gln Pro Cys Cys  
 65 70 75 80  
 Arg Pro Thr Arg Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser Thr  
 85 90 95  
 Trp Arg Thr Val Asp Arg Leu Ser Ala Thr Ala Cys Gly Cys Leu Gly  
 100 105 110

<210> 9  
 <211> 111  
 <212> PRT  
 <213> Homo sapiens

<400> 9  
 Gly Pro Gly Ser Arg Ala Arg Ala Ala Gly Ala Arg Gly Cys Arg Leu  
 1 5 10 15  
 Arg Ser Gln Leu Val Pro Val Arg Ala Leu Gly Leu Gly His Arg Ser

20	25	30
Asp Glu Leu Val Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg Arg Ala		
35	40	45
Arg Ser Pro His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala Gly Ala		
50	55	60
Leu Arg Pro Pro Pro Gly Ser Arg Pro Val Ser Gln Pro Cys Cys Arg		
65	70	75
Pro Thr Arg Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser Thr Trp		
85	90	95
Arg Thr Val Asp Arg Leu Ser Ala Thr Ala Cys Gly Cys Leu Gly		
100	105	110

<210> 10  
 <211> 110  
 <212> PRT  
 <213> Homo sapiens

<400> 10
Pro Gly Ser Arg Ala Arg Ala Ala Gly Ala Arg Gly Cys Arg Leu Arg
1 5 10 15
Ser Gln Leu Val Pro Val Arg Ala Leu Gly Leu Gly His Arg Ser Asp
20 25 30
Glu Leu Val Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg Arg Ala Arg
35 40 45
Ser Pro His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala Gly Ala Leu
50 55 60
Arg Pro Pro Pro Gly Ser Arg Pro Val Ser Gln Pro Cys Cys Arg Pro
65 70 75 80
Thr Arg Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser Thr Trp Arg
85 90 95
Thr Val Asp Arg Leu Ser Ala Thr Ala Cys Gly Cys Leu Gly
100 105 110

<210> 11  
 <211> 109  
 <212> PRT  
 <213> Homo sapiens

<400> 11
Gly Ser Arg Ala Arg Ala Ala Gly Ala Arg Gly Cys Arg Leu Arg Ser
1 5 10 15
Gln Leu Val Pro Val Arg Ala Leu Gly Leu Gly His Arg Ser Asp Glu
20 25 30
Leu Val Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg Arg Ala Arg Ser
35 40 45
Pro His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala Gly Ala Leu Arg
50 55 60
Pro Pro Pro Gly Ser Arg Pro Val Ser Gln Pro Cys Cys Arg Pro Thr
65 70 75 80
Arg Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser Thr Trp Arg Thr
85 90 95
Val Asp Arg Leu Ser Ala Thr Ala Cys Gly Cys Leu Gly
100 105

<210> 12  
 <211> 108  
 <212> PRT

<213> Homo sapiens

<400> 12

Ser Arg Ala Arg Ala Ala Gly Ala Arg Gly Cys Arg Leu Arg Ser Gln  
1 5 10 15  
Leu Val Pro Val Arg Ala Leu Gly Leu Gly His Arg Ser Asp Glu Leu  
20 25 30  
Val Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg Arg Ala Arg Ser Pro  
35 40 45  
His Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala Gly Ala Leu Arg Pro  
50 55 60  
Pro Pro Gly Ser Arg Pro Val Ser Gln Pro Cys Cys Arg Pro Thr Arg  
65 70 75 80  
Tyr Glu Ala Val Ser Phe Met Asp Val Asn Ser Thr Trp Arg Thr Val  
85 90 95  
Asp Arg Leu Ser Ala Thr Ala Cys Gly Cys Leu Gly  
100 105

<210> 13

<211> 107

<212> PRT

<213> Homo sapiens

<400> 13

Arg Ala Arg Ala Ala Gly Ala Arg Gly Cys Arg Leu Arg Ser Gln Leu  
1 5 10 15  
Val Pro Val Arg Ala Leu Gly Leu Gly His Arg Ser Asp Glu Leu Val  
20 25 30  
Arg Phe Arg Phe Cys Ser Gly Ser Cys Arg Arg Ala Arg Ser Pro His  
35 40 45  
Asp Leu Ser Leu Ala Ser Leu Leu Gly Ala Gly Ala Leu Arg Pro Pro  
50 55 60  
Pro Gly Ser Arg Pro Val Ser Gln Pro Cys Cys Arg Pro Thr Arg Tyr  
65 70 75 80  
Glu Ala Val Ser Phe Met Asp Val Asn Ser Thr Trp Arg Thr Val Asp  
85 90 95  
Arg Leu Ser Ala Thr Ala Cys Gly Cys Leu Gly  
100 105

<210> 14

<211> 106

<212> PRT

<213> Homo sapiens

<400> 14

Ala Arg Ala Ala Gly Ala Arg Gly Cys Arg Leu Arg Ser Gln Leu Val  
1 5 10 15  
Pro Val Arg Ala Leu Gly Leu Gly His Arg Ser Asp Glu Leu Val Arg  
20 25 30  
Phe Arg Phe Cys Ser Gly Ser Cys Arg Arg Ala Arg Ser Pro His Asp  
35 40 45  
Leu Ser Leu Ala Ser Leu Leu Gly Ala Gly Ala Leu Arg Pro Pro Pro  
50 55 60  
Gly Ser Arg Pro Val Ser Gln Pro Cys Cys Arg Pro Thr Arg Tyr Glu  
65 70 75 80  
Ala Val Ser Phe Met Asp Val Asn Ser Thr Trp Arg Thr Val Asp Arg  
85 90 95

Leu Ser Ala Thr Ala Cys Gly Cys Leu Gly  
100 105

<210> 15  
<211> 105  
<212> PRT  
<213> Homo sapiens

<400> 15  
Arg Ala Ala Gly Ala Arg Gly Cys Arg Leu Arg Ser Gln Leu Val Pro  
1 5 10 15  
Val Arg Ala Leu Gly Leu Gly His Arg Ser Asp Glu Leu Val Arg Phe  
20 25 30  
Arg Phe Cys Ser Gly Ser Cys Arg Arg Ala Arg Ser Pro His Asp Leu  
35 40 45  
Ser Leu Ala Ser Leu Leu Gly Ala Gly Ala Leu Arg Pro Pro Pro Gly  
50 55 60  
Ser Arg Pro Val Ser Gln Pro Cys Cys Arg Pro Thr Arg Tyr Glu Ala  
65 70 75 80  
Val Ser Phe Met Asp Val Asn Ser Thr Trp Arg Thr Val Asp Arg Leu  
85 90 95  
Ser Ala Thr Ala Cys Gly Cys Leu Gly  
100 105

<210> 16  
<211> 104  
<212> PRT  
<213> Homo sapiens

<400> 16  
Ala Ala Gly Ala Arg Gly Cys Arg Leu Arg Ser Gln Leu Val Pro Val  
1 5 10 15  
Arg Ala Leu Gly Leu Gly His Arg Ser Asp Glu Leu Val Arg Phe Arg  
20 25 30  
Phe Cys Ser Gly Ser Cys Arg Arg Ala Arg Ser Pro His Asp Leu Ser  
35 40 45  
Leu Ala Ser Leu Leu Gly Ala Gly Ala Leu Arg Pro Pro Pro Gly Ser  
50 55 60  
Arg Pro Val Ser Gln Pro Cys Cys Arg Pro Thr Arg Tyr Glu Ala Val  
65 70 75 80  
Ser Phe Met Asp Val Asn Ser Thr Trp Arg Thr Val Asp Arg Leu Ser  
85 90 95  
Ala Thr Ala Cys Gly Cys Leu Gly  
100

<210> 17  
<211> 103  
<212> PRT  
<213> Homo sapiens

<400> 17  
Ala Gly Ala Arg Gly Cys Arg Leu Arg Ser Gln Leu Val Pro Val Arg  
1 5 10 15  
Ala Leu Gly Leu Gly His Arg Ser Asp Glu Leu Val Arg Phe Arg Phe  
20 25 30  
Cys Ser Gly Ser Cys Arg Arg Ala Arg Ser Pro His Asp Leu Ser Leu  
35 40 45

Ala Ser Leu Leu Gly Ala Gly Ala Leu Arg Pro Pro Pro Gly Ser Arg  
 50 55 60  
 Pro Val Ser Gln Pro Cys Cys Arg Pro Thr Arg Tyr Glu Ala Val Ser  
 65 70 75 80  
 Phe Met Asp Val Asn Ser Thr Trp Arg Thr Val Asp Arg Leu Ser Ala  
 85 90 95  
 Thr Ala Cys Gly Cys Leu Gly  
 100

<210> 18

<211> 102

<212> PRT

<213> Homo sapiens

<400> 18

Gly Ala Arg Gly Cys Arg Leu Arg Ser Gln Leu Val Pro Val Arg Ala  
 1 5 10 15  
 Leu Gly Leu Gly His Arg Ser Asp Glu Leu Val Arg Phe Arg Phe Cys  
 20 25 30  
 Ser Gly Ser Cys Arg Arg Ala Arg Ser Pro His Asp Leu